RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/555,094Source: PCTDate Processed by STIC: 11/88/2005

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 11/08/2005
PATENT APPLICATION: US/10/555,074 TIME: 12:52:31

Input Set : A:\seqlisting.txt

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2 <110> APPLICANT: JAPAN SCIENCE AND TECHNOLOGY AGENCY
      4 <120> TITLE OF INVENTION: Human antibody against human interleukin-18, the
              antibody fragment, and the use thereof
      7 <130> FILE REFERENCE: A211-04PCT
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/555,074
C--> 9 <141> CURRENT FILING DATE: 2005-10-28
      9 <150> PRIOR APPLICATION NUMBER: JP 2003-125948
     10 <151> PRIOR FILING DATE: 2003-04-30
     12 <160> NUMBER OF SEQ ID NOS: 12
     14 <170> SOFTWARE: PatentIn Ver. 2.1
     16 <210> SEQ ID NO: 1
     17 <211> LENGTH: 357
     18 <212> TYPE: DNA
     19 <213> ORGANISM: Homo sapiens
     21 <400> SEQUENCE: 1
     22 caggtgcagc tggtgcagtc tggggctgag gtgaggaggc ctgggggcctc agtgagggtt 60
     23 teetgeaagg catetggata cacetteact agteactata tacaetgggt gegacaggee 120
     24 cctggacaag ggcttgagtg ggtggcaata atcaacccta gtgatggcag aacagactac 180
     25 gcacagaagt tccagggcag agtcaccgtg accagggaca cgtccgcgag cagtgtctac 240
     26 atgggaataa gcagcctgag atctgaggac acggccatgt attactgtgc gagaacagcg 300
     27 cgtggattca gttatgcgac agactggggc cagggaaccc tggtcaccgt ctcctca
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     30 <210> SEO ID NO: 2
     31 <211> LENGTH: 357
     32 <212> TYPE: DNA
     33 <213> ORGANISM: Homo sapiens
     35 <220> FEATURE:
     36 <221> NAME/KEY: CDS
     37 <222> LOCATION: (1)..(357)
     39 <400> SEQUENCE: 2
     40 cag gtg cag ctg gtg cag tct ggg gct gag gtg agg agg cct ggg gcc
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     41 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Arg Arg Pro Gly Ala
     42
          1
                                                                  15
     44 tca gtg agg gtt tcc tgc aag gca tct gga tac acc ttc act agt cac
                                                                           96
     45 Ser Val Arg Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser His
     46
                     20
                                         25
                                                              30
    48 tat ata cac tgg gtg cga cag gcc cct gga caa ggg ctt gag tgg gtg
                                                                           144
    49 Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Val
     50
                 35
    52 gca ata atc aac cct agt gat ggc aga aca gac tac gca cag aag ttc
                                                                           192
    53 Ala Ile Ile Asn Pro Ser Asp Gly Arg Thr Asp Tyr Ala Gln Lys Phe
             50
     54
                                 55
                                                      60
    56 cag ggc aga gtc acc gtg acc agg gac acg tcc gcg agc agt gtc tac
                                                                           240
    57 Gln Gly Arg Val Thr Val Thr Arg Asp Thr Ser Ala Ser Ser Val Tyr
```

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75
58 65
                         70
                                                                   80
60 atg gga ata agc agc ctg aga tct gag gac acg gcc atg tat tac tgt
                                                                        288
61 Met Gly Ile Ser Ser Leu Arg Ser Glu Asp Thr Ala Met Tyr Tyr Cys
62
                     85
64 gcg aga aca gcg cgt gga ttc agt tat gcg aca gac tgg ggc cag gga
                                                                        336
65 Ala Arg Thr Ala Arg Gly Phe Ser Tyr Ala Thr Asp Trp Gly Gln Gly
66
               100
                                    105
                                                         110
                                                                        357
68 acc ctg gtc acc gtc tcc tca
69 Thr Leu Val Thr Val Ser Ser
70
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73 <210> SEQ ID NO: 3
74 <211> LENGTH: 119
75 <212> TYPE: PRT
76 <213> ORGANISM: Homo sapiens
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79 Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Arg Arg Pro Gly Ala
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                                                              15
                                         10
82 Ser Val Arg Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Ser His
                20
83
                                     25
85 Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Val
86
            35
88 Ala Ile Ile Asn Pro Ser Asp Gly Arg Thr Asp Tyr Ala Gln Lys Phe
        50
89
                             55
                                                  60
91 Gln Gly Arg Val Thr Val Thr Arg Asp Thr Ser Ala Ser Ser Val Tyr
92 65
                         70
                                              75
                                                                   80
94 Met Gly Ile Ser Ser Leu Arg Ser Glu Asp Thr Ala Met Tyr Tyr Cys
95
                     85
97 Ala Arg Thr Ala Arg Gly Phe Ser Tyr Ala Thr Asp Trp Gly Gln Gly
               100
98
                                    105
                                                         110
100 Thr Leu Val Thr Val Ser Ser
101
            115
105 <210> SEQ ID NO: 4
106 <211> LENGTH: 5
107 <212> TYPE: PRT
108 <213 > ORGANISM: Homo sapiens
110 <400> SEQUENCE: 4
111 Ser His Tyr Ile His
112
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115 <210> SEQ ID NO: 5
116 <211> LENGTH: 17
117 <212> TYPE: PRT
118 <213> ORGANISM: Homo sapiens
120 <400> SEQUENCE: 5
121 Ile Ile Asn Pro Ser Asp Gly Arg Thr Asp Tyr Ala Gln Lys Phe Gln
122
                                                               15
      1
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                                          10
124 Gly
128 <210> SEQ ID NO: 6
129 <211> LENGTH: 10
130 <212> TYPE: PRT
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133 <400> SEQUENCE: 6
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138 <210> SEQ ID NO: 7
139 <211> LENGTH: 327
140 <212> TYPE: DNA
141 <213> ORGANISM: Homo sapiens
143 <400> SEQUENCE: 7
144 tectatgage tgaeteagee acceteggtg teagtgteee eaggaeaaae ggeeaggate 60
145 acctgctctg gagatgcatt gccaaaaaaa tatgcttatt ggtaccagca gaagccaggc 120
146 caggcccctg tgctggtgat atataaagac agtgagaggc cctcagggat ccctgagcga 180
147 ttctctggct ccagctcagg gacaacagtc acgttgacca tcagtggagt ccaggcagaa 240
148 gacgaggetg actattactg tcaatcagca gacagcagtg gtacttatgt ggtattegge 300
                                                                        327
149 ggagggaccc agctcaccgt tttaggt
152 <210> SEQ ID NO: 8
153 <211> LENGTH: 327
154 <212> TYPE: DNA
.155 <213> ORGANISM: Homo sapiens
157 <220> FEATURE:
158 <221> NAME/KEY: CDS
159 <222> LOCATION: (1)..(327)
161 <400> SEQUENCE: 8
162 tcc tat gag ctg act cag cca ccc tcg gtg tca gtg tcc cca gga caa
                                                                        48
163 Ser Tyr Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
                       5
                                          10
                                                               15
164
      1
166 acg gcc agg atc acc tgc tct gga gat gca ttg cca aaa aaa tat gct
                                                                        96
167 Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Lys Tyr Ala
                                                           30
168
                  20
                                      25
170 tat tog tac cag cag aag cca ggc cag gcc cct gtg ctg gtg ata tat
                                                                        144
171 Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
                                                       45
172
             35
                                  40
.174 aaa gac agt gag agg ccc tca ggg atc cct gag cga ttc tct ggc tcc
                                                                        192
175 Lys Asp Ser Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
176
         50
                              55
                                                  60
178 agc tca ggg aca aca gtc acg ttg acc atc agt gga gtc cag gca gaa
                                                                        240
179 Ser Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
180 65
                                                                   80
                          70
182 gac gag gct gac tat tac tgt caa tca gca gac agc agt ggt act tat
                                                                        288
183 Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Thr Tyr
184
                      85
                                          90
                                                               95
                                                                        327
186 gtg gta ttc ggc gga ggg acc cag ctc acc gtt tta ggt
187 Val Val Phe Gly Gly Gly Thr Gln Leu Thr Val Leu Gly
188
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191 <210> SEQ ID NO: 9
192 <211> LENGTH: 109
193 <212> TYPE: PRT
194 <213> ORGANISM: Homo sapiens
196 <400> SEQUENCE: 9
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PATENT APPLICATION: US/10/555,074 TIME: 12:52:31

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197 Ser Tyr Glu Leu Thr Gln Pro Pro Ser Val Ser Val Ser Pro Gly Gln
                                          10
                                                               15
198
      1
200 Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Lys Tyr Ala
201
203 Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
204
             35
                                  40
206 Lys Asp Ser Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
         50
                              55
207
209 Ser Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
                         70
                                              75
                                                                   80
210 65
212 Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Thr Tyr
                                          90
213
                                                               95
215 Val Val Phe Gly Gly Gly Thr Gln Leu Thr Val Leu Gly
216
                100
                                     105
220 <210> SEQ ID NO: 10
221 <211> LENGTH: 11
222 <212> TYPE: PRT
223 <213> ORGANISM: Homo sapiens
225 <400> SEQUENCE: 10
226 Ser Gly Asp Ala Leu Pro Lys Lys Tyr Ala Tyr
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                                          10
230 <210> SEQ ID NO: 11
231 <211> LENGTH: 7
232 <212> TYPE: PRT
233 <213> ORGANISM: Homo sapiens
235 <400> SEQUENCE: 11
236 Lys Asp Ser Glu Arg Pro Ser
237
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240 <210> SEQ ID NO: 12
241 <211> LENGTH: 11
242 <212> TYPE: PRT
243 <213> ORGANISM: Homo sapiens
245 <400> SEQUENCE: 12
246 Gln Ser Ala Asp Ser Ser Gly Thr Tyr Val Val
247
                                          10
      1/10
251
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VERIFICATION SUMMARYDATE: 11/08/2005PATENT APPLICATION: US/10/555,074TIME: 12:52:32

Input Set : A:\seqlisting.txt

Output Set: N:\CRF4\11082005\J555074.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No

L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date